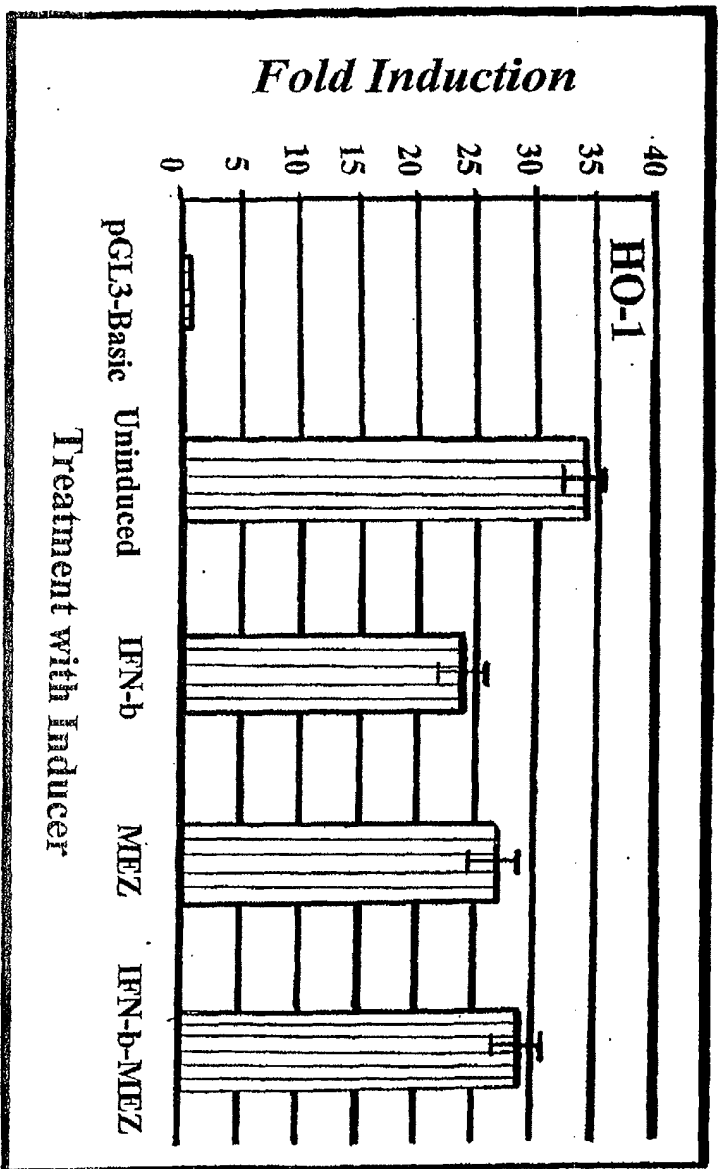


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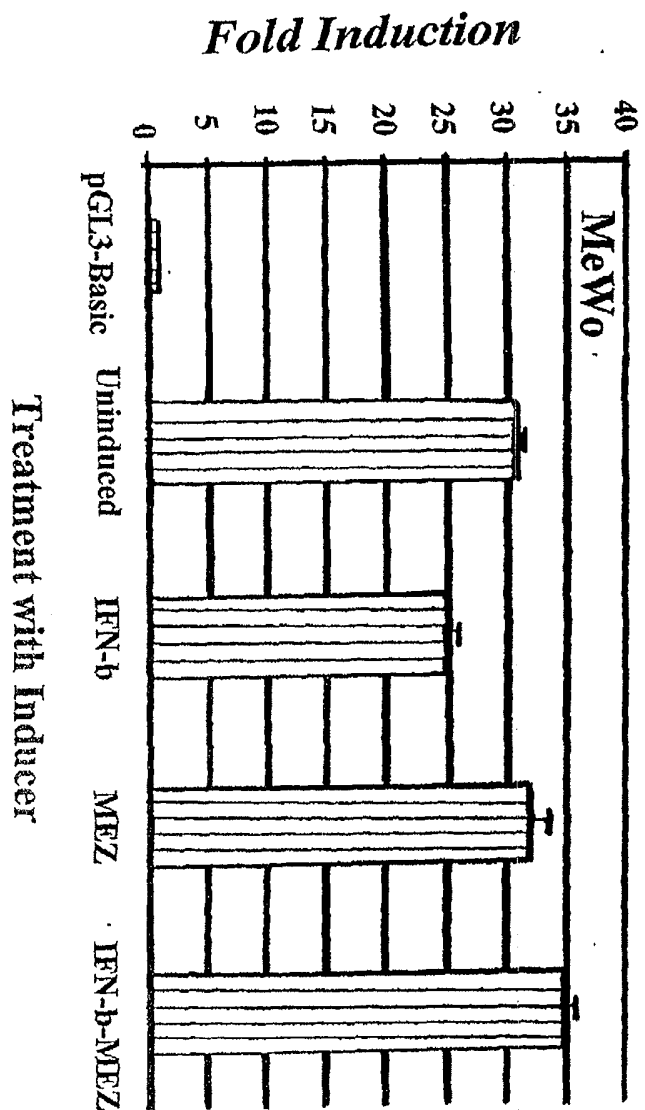
FIGURE 1A

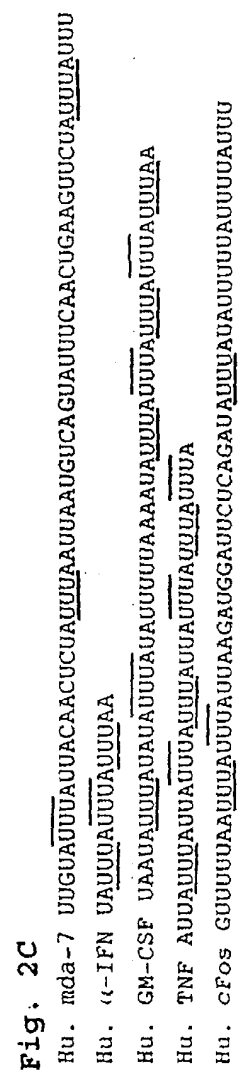
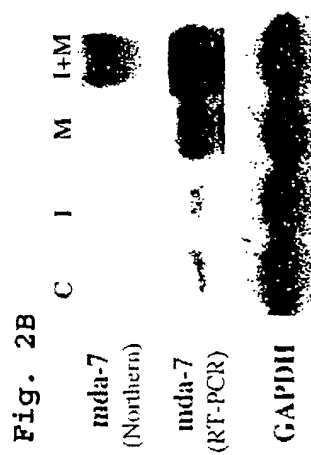
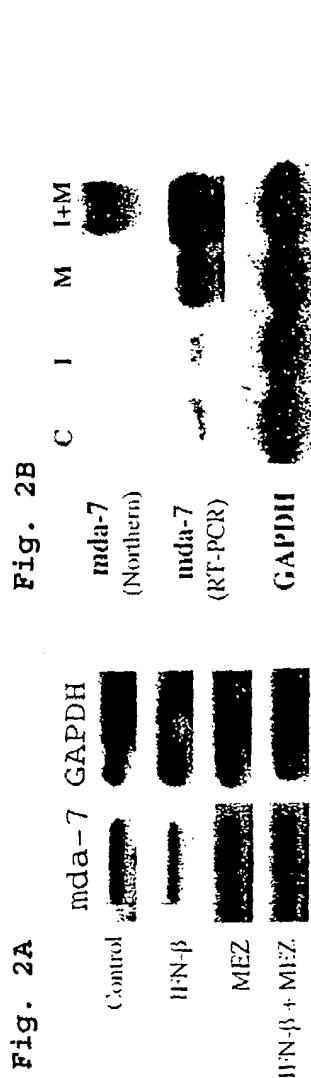


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FIGURE 1B





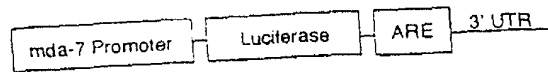


Fig. 3A

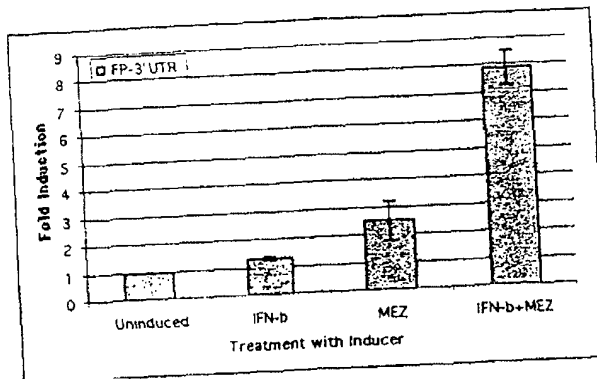


Fig. 3B

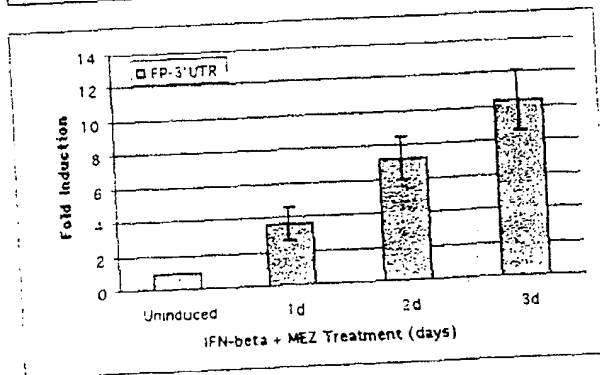


Fig. 4A

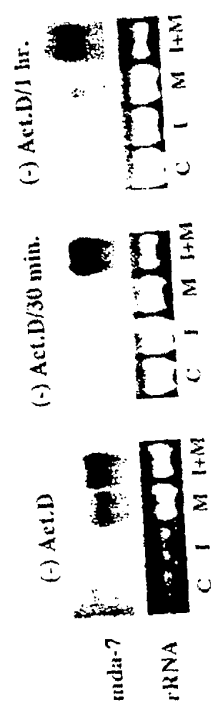


Fig. 4B

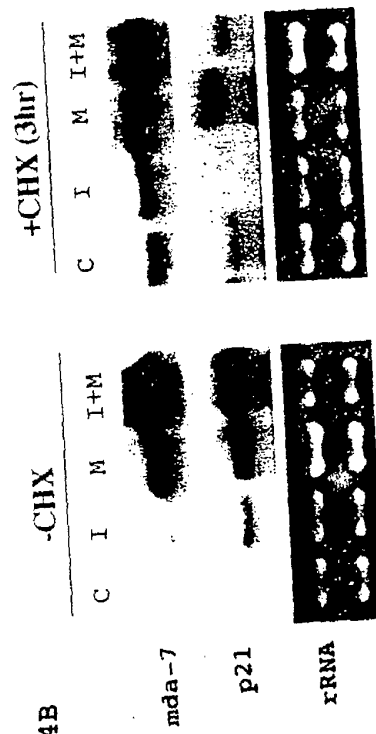


Fig. 5A

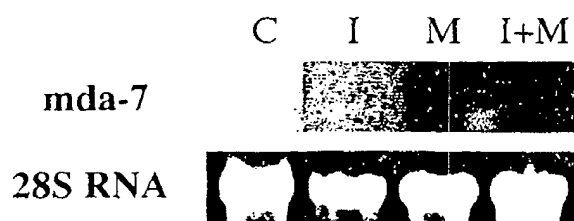


Fig. 5B

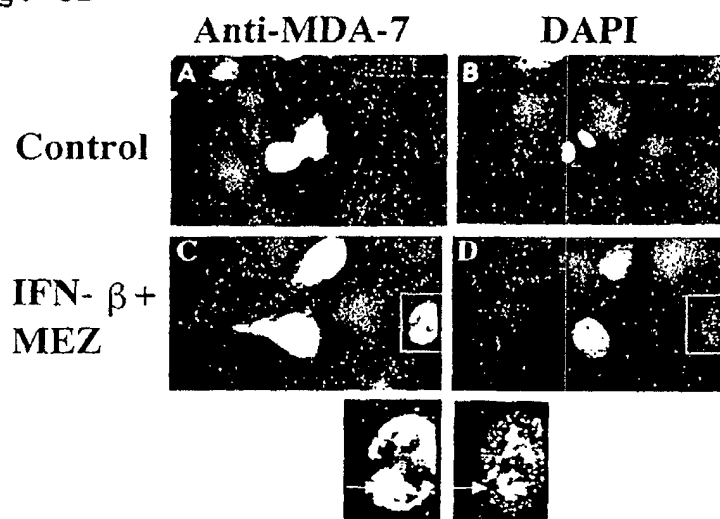
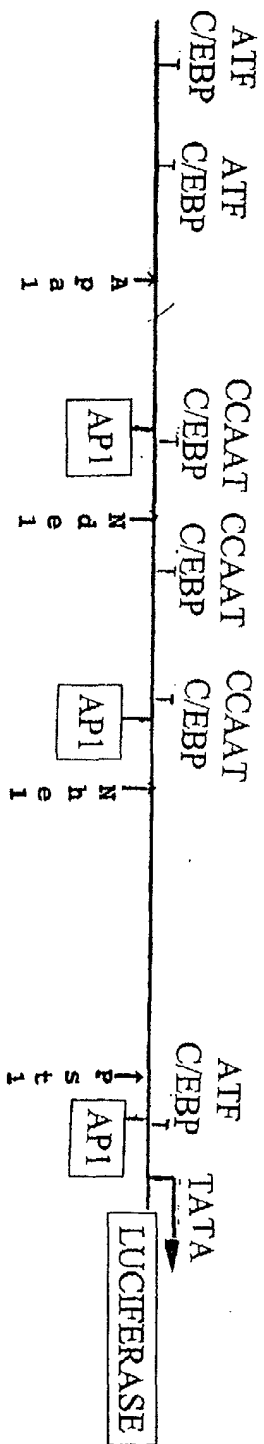


Fig. 6

TAATACGACTCACTATAGGGCGTCGACTCGATCACCTTTTGAACCCAGGTCTGCCTGCCTCCAAAGCTTGACTCATAAC -2161
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 CTCATCGGAGGCTGAGGCCCAGGCACATGTTGCCTGAAGTATCCATGTTATATGATTCTTCCTCAGACAGAGTGAGCT -2001
 ACTCACGATCCCAGGTGTACCCTGAGGCCAGCCAAGGTGTATCCATGACCTCATGCCTCTGTTCCAGCCTGCCCTTTAAC -1921
 AGCTCATCCCACCTGCCTGCCCTCCCCGCCTATCTGCAGACAGTAGTCTAGGATTTTCAGCTGCCTGGGGGCTCATTTTC -1841
 CCTCTCAGCTTCCTGCTTTAGCTGTCTCCTGCCTCCCACTCACCTATTACTCCAGCACTCTCACCTGGTCTCTTTTCTG -1761
 TCTCATCACTGCCTCTTGACATCTTTATCTCATAGTAGTTAGTTAGGGGTTCTTGGTAATGCCCTAAATCCACATGGTGG -1681
 GAAGGGGGGAGTGGGGGAAGAGAGTGCCTGTGGGGCTGTGCCTACTTCTGGAGGGTAAGACTCGGGCCCTCCAGGAACA -1601
 AAGGATTCAGGCTGGTGGCAGCTATAGCCAAGCAGACTGCTGGCCAGGGATTGCAAAGGAGTATTTGTTTGCTTAAGAA -1521
 AATAACAACACTGAGTATGAGATGGAGGGAGGGGGTGTGGTGCCAGAGAGATTGGGAAGAGTCTGCCAAGGGTGTGTT -1441
 CTACTCACTCTCTCTTTCTTTTCTCTCCACTGAGCTGGAGGCAGTTATCTGTCCCCCACGTCACATTCCTACTCCCG -1361
 AP1
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 ATGGTCTCAGTCACAACACTACTCATCTCTGCCTCTGTAGCACGAAAGCAATTAGCAACAATATGTCAACAAACATATGTGA -1201
 CCCCATGAAAACCTTTATTTATTATGGATACGGAACCTGAAAATAATGTCTTTCTTTTGATTTTTCCTCAATCATTAAA -1121
 C/EBP
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 ACAGTTGGCCAATCCCTGATTCCCAAAATGTATTCCTCAGGGATGTGGGCAAATACTTATGGGAAGTGTGGATTAAACA -961
 AP1 C/EBP
 GAGTTAAGAAGCATCAGACATTTCCAGGACGGGCTAGCACATGCCAGGGCTCTCTAACTGACCTCATTGGATTCTCTGT -881
 TTCATGGAGGATCTTGCAAGACAAGAAATTCCTCAAACCTAGAGTCTGAGGACTGTGCTTTGGGAAACACTGCTCTGCTTG -801
 ATGCCCTCACTGGGCACATGGTAGAATCTAGAGCTGAGTGCCTTGCTAGCTGGAGATAGGGTCAGAGCTCTTGACTGCCC -721
 TGGCAGTCTTGACACATCACGCTGTCTGTGTCCCTGAGTGGTTCAGAGCCACACAGGCCAAGACTAGCCCACCAGAGCA -641
 AP1
 CCAGGCTCCAGCTTTCTGGGCTTGTCATGCGTACATTTCTTATTCTTCTGCTTTCCAGAACCTAAGGAGAGGCAC -561
 ATTTTGGTTGAGTGATTATAACCTAGGGACCATGGGTAGCTGCATGTGAGGAAACACTCTCAACTTCTGGCCCTGAT -481
 GGATTAAAGGAGAGGTACTTACAGGTTATTTCTTCGCTGTGGACTACTGTCCAGCATGAATAGGGCATCATTATTGAAT -401
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 TAAGAGACTTGCCCTGCCCACACAGCCAGTGGTAGAGCCTGAACTCAAACCCAGGTCTCATCTCACCTCAGGGGCTGCT -161
 TTCCCATCGCTGTATTGTCCTTAAAGTGATGGGTGACTAGGCAATGAAGTAATTCTCTAGGAAAGCATGACCAATTTCC -81
 AP1 C/EBP
 CTTTCTCCACCTCCCTCTTTTCTCCACCCCTCCCCCATCAGCCCCCATATATATGCCCAAATCTCCACAAAGCCTTGC 0
 TTGCCTGCAAACCTTTACTTCTGAAATGACTTCCACGGCTGGGACG +45

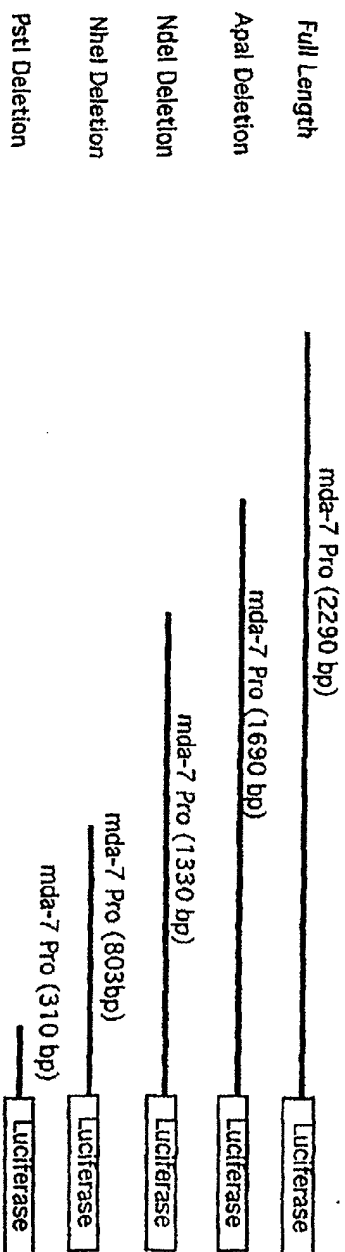
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FIGURE 7A



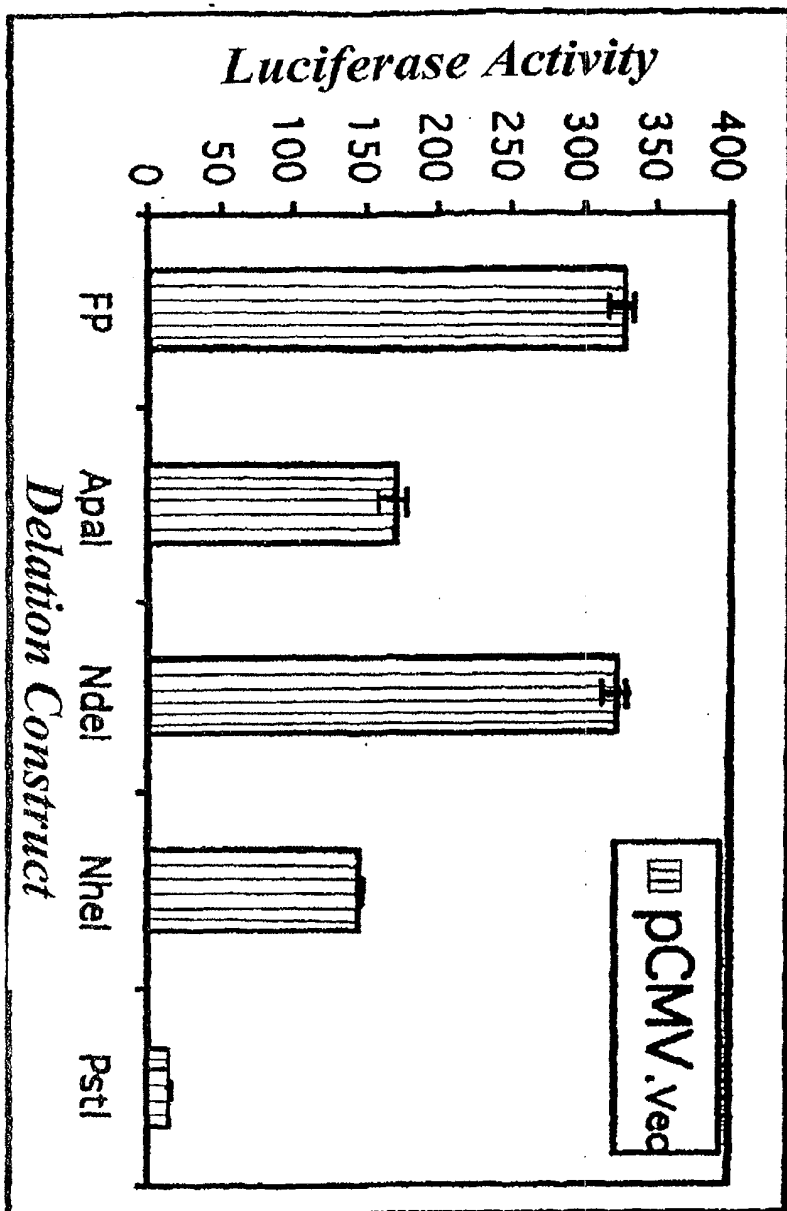
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FIGURE 7B



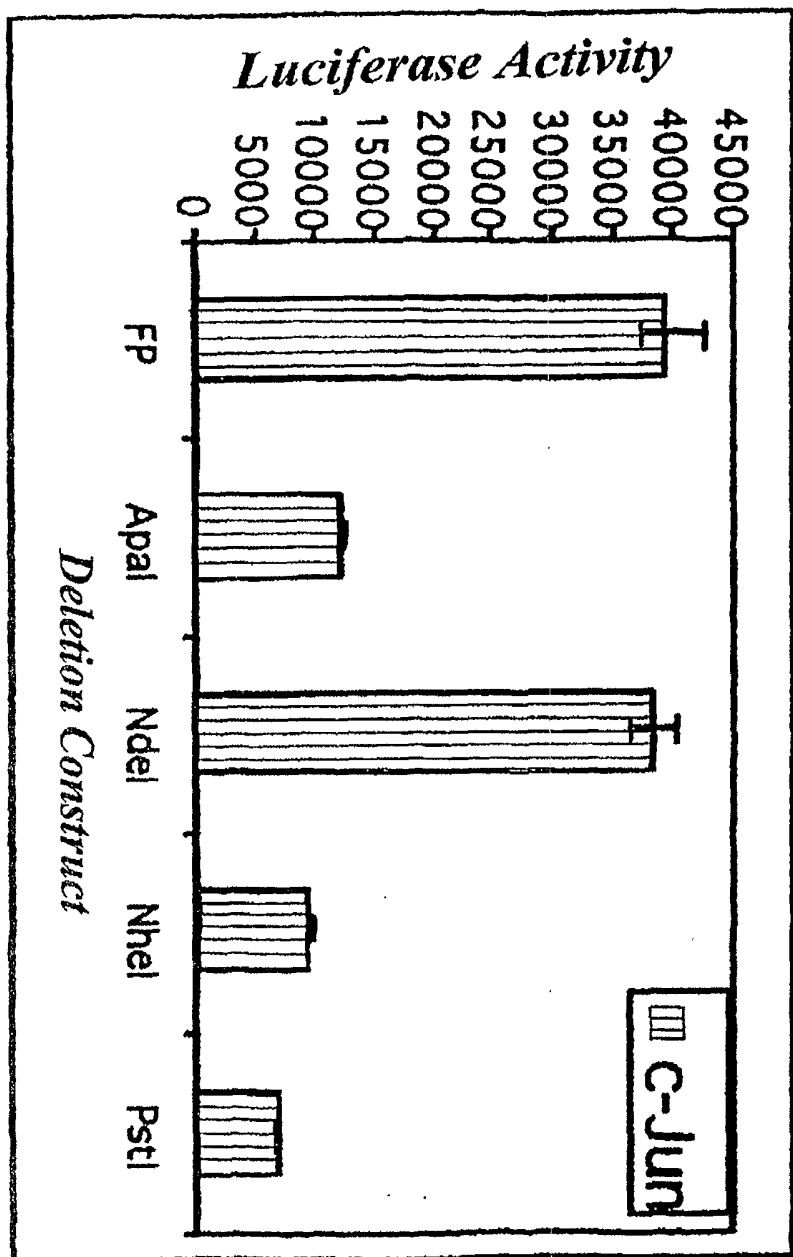
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FIGURE 7C



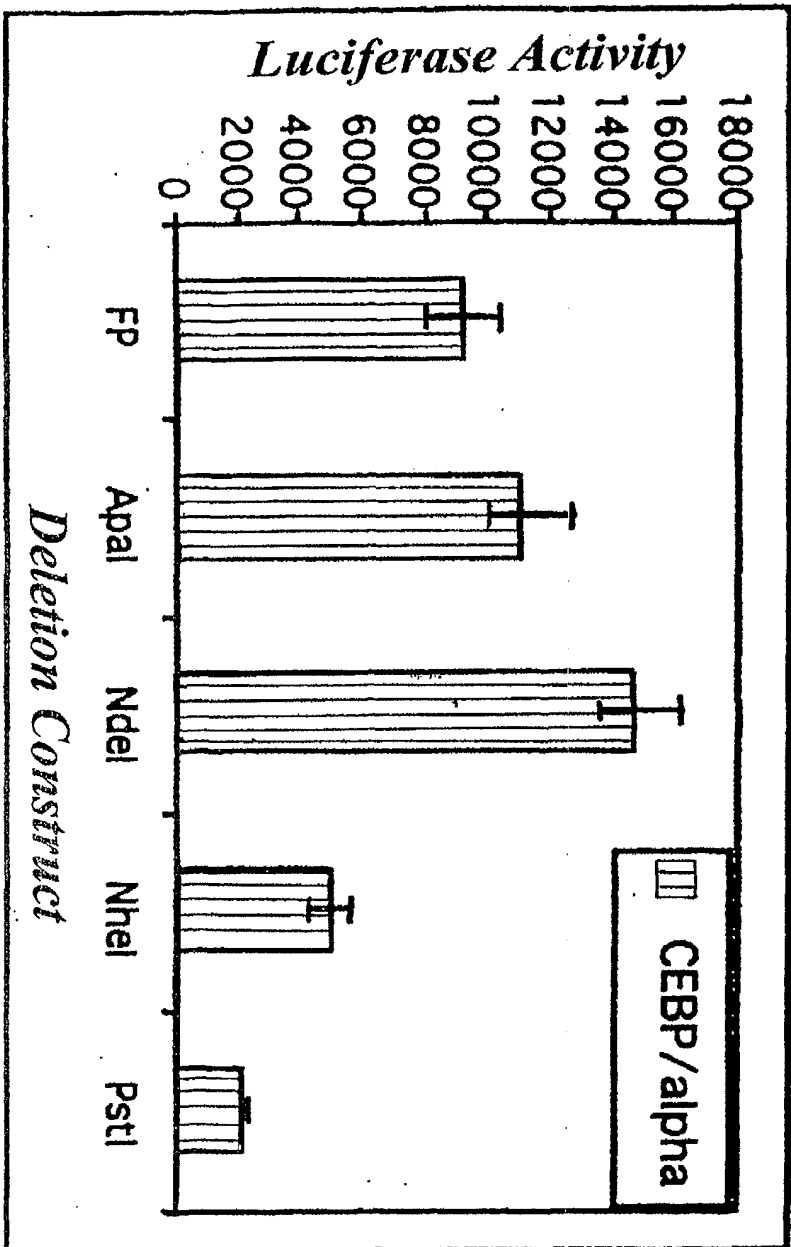
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FIGURE 7D



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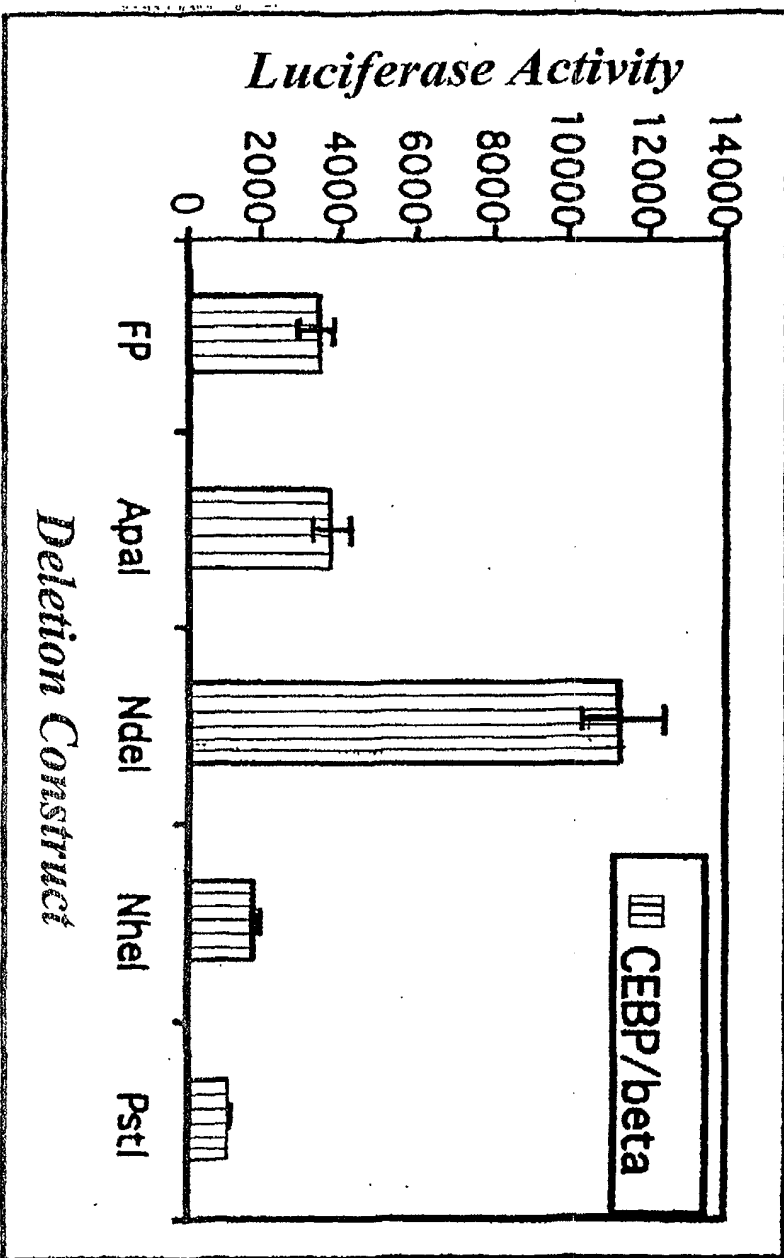
FIGURE 7E



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FIGURE 7F

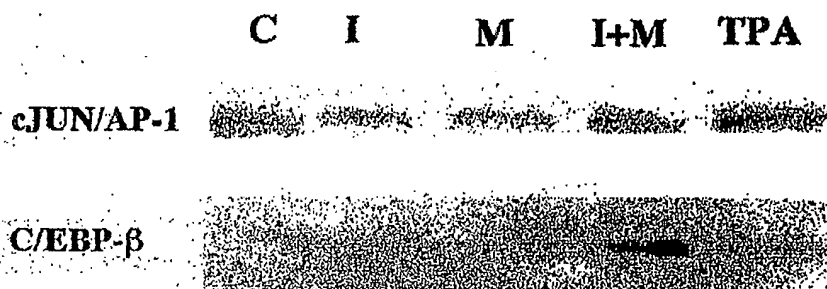


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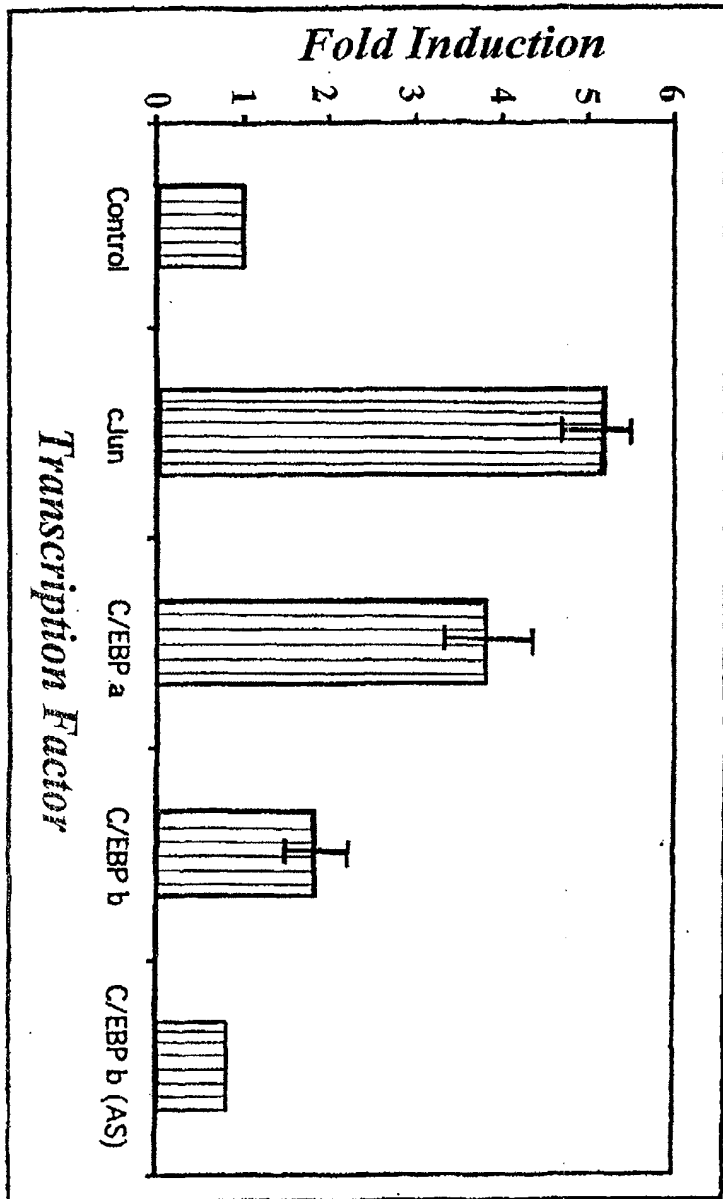
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FIGURE 8A



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FIGURE 8B



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FIGURE 8C

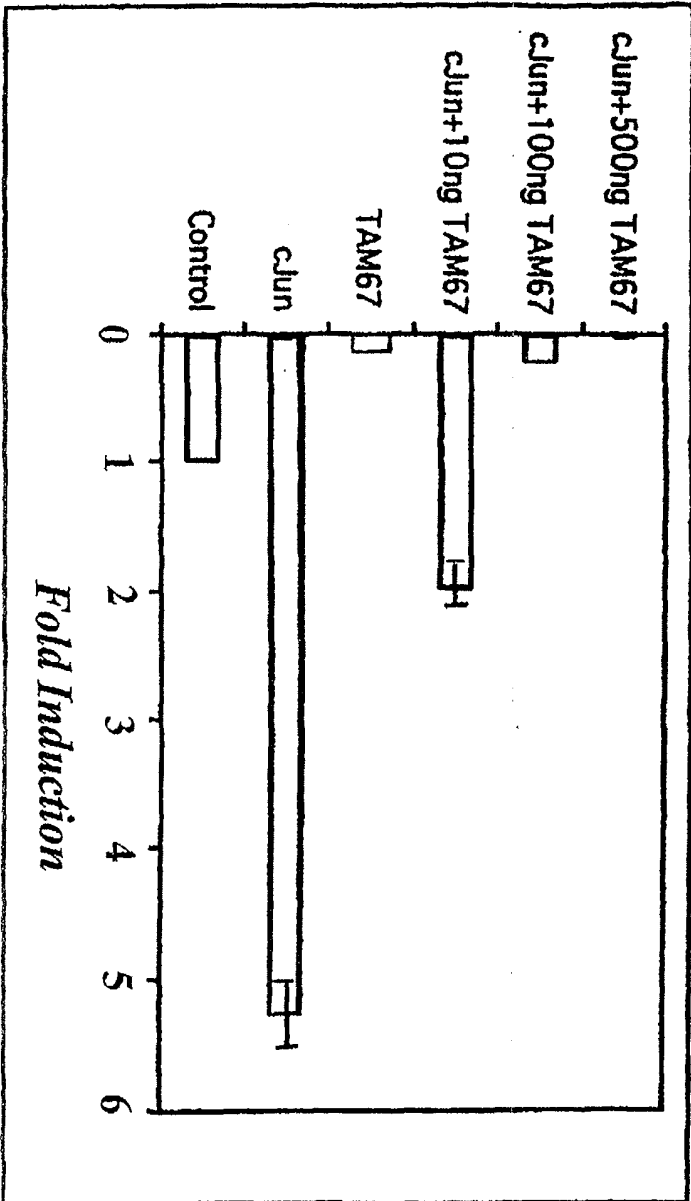


Fig. 9A

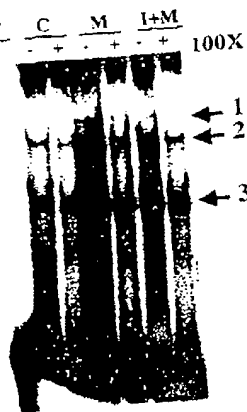


Fig. 9C

C/EBP-β
WT 10X 100X MT

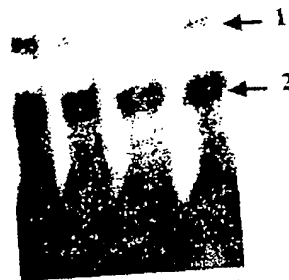


Fig. 9E

E
Protein - + + +
Anti-CEBP-β - - 1μg 5μg

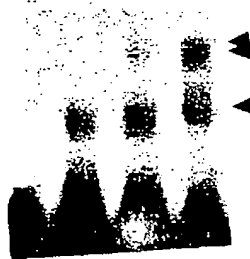
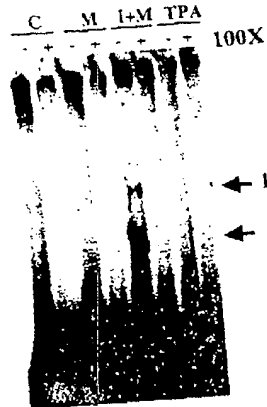
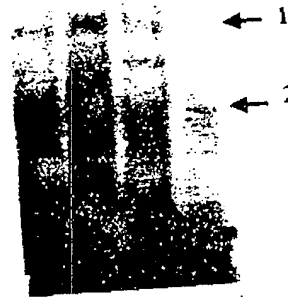


Fig. 9B



cJUN/AP-1
WT MT 10X 100X

Fig. 9D



Protein
Anti-AP-1
- + + +
- - 1μg 5μg

Fig. 9F

